

Web Images Maps News Shopping Gmail more ▼



Sign in

compare source article using alignment algorithm to compute edit distance

Search


Advanced Search  
Preferences



Web: Result details: [SIMIL](#)

Results 1 - 10 of about 19,100 for [compare source article using alignment algorithm to compute edit distance](#). (0.23 seconds)

## Scholarly articles for [compare source article using alignment algorithm to compute edit distance](#)

-  [A comparison of alignment models for statistical machine ...](#) - Och - Cited by 94
- [Unsupervised construction of large paraphrase corpora: ...](#) - Dolan - Cited by 60
- [Practical algorithms for transposition-invariant ...](#) - Lemström - Cited by 8

### 1. [Levenshtein distance](#) - Wikipedia, the free encyclopedia

This **algorithm** is based on the Wagner-Fischer **algorithm** for **edit distance**. .... that are useful in applications which **compute** many of them and **compare** them. ...  
[en.wikipedia.org/wiki/Levenshtein\\_distance](#) - 41k - [Cached](#) - [Similar pages](#)

### 2. [Edit distance-based kernel functions for structural pattern ...](#)

[2] Special Section on Graph **Algorithms** and **Computer Vision**, IEEE Trans. .... E. , Computing approximate **tree edit distance** using relaxation labeling. ...  
[portal.acm.org/citation.cfm?id=1221253](#) - [Similar pages](#)  
by M Neuhäus - 2006 - [Cited by 3](#) - [Related articles](#) - [All 3 versions](#)

### 3. [Edit distance between two RNA structures](#)

Several **algorithms** are presented to **compute** the **edit distance** two RNA sequences with ... RNAling program: **alignment** of RNA sequences **using** both primary and ...  
[portal.acm.org/citation.cfm?id=369133.369214](#) - [Similar pages](#)  
by GH Lin - 2001 - [Cited by 7](#) - [Related articles](#) - [All 8 versions](#)  
[More results from portal.acm.org ...](#)

### 4. [Biomolecular Engineering : Peak alignment using restricted edit ...](#)

Here the intermediate values  $d_{ij}$  denote the **edit distance** of .... as we use the same technique later on to **compute distance** Click to view the MathML **source** ...  
[linkinghub.elsevier.com/retrieve/pii/S138903440700024X](#) - [Similar pages](#)  
by V Mäkinen - 2007 - [Cited by 1](#) - [Related articles](#) - [All 5 versions](#)

### 5. [IngentaConnect Fast, Optimal Alignment of Three Sequences Using ...](#)

**Alignment algorithms** can be used to infer a relationship between sequences when the ... It is particularly fast when the (three-way) **edit distance** is small. ...  
[www.ingentaconnect.com/content/ap/jv/2000/00000207/00000003/art02177](#)  
- [Similar pages](#)  
by DR POWELL - 2000 - [Cited by 14](#) - [Related articles](#) - [All 4 versions](#)

### 6. [LNCS 4707 - An Efficient Sequence Alignment Algorithm on a LARPBS](#)

become a major field as many **articles** have been devoted to it. It consists in **using**. **computer science** for designing efficient **algorithms** in order to solve ...  
[www.springerlink.com/index/e404581214864231.pdf](#) - [Similar pages](#)

by D Somé - [Related articles](#)

7. [\[PDF\] Similarity of names across scripts: Edit distance using learned ...](#)

File Format: PDF/Adobe Acrobat - [View as HTML](#)

We **compare** two strings **using** an **algorithm** that .... Performance: This new way to **compute edit distance** has a higher complexity. In the- ...  
[langtech.jrc.it/Documents/2008\\_Gotal\\_Pouliquen\\_edit-distance.pdf](#) - [Similar pages](#)  
by B Pouliquen

8. [SUSAX: Context-Specific Searching in XML Documents Using Sequence ...](#)

**Source** Paths. are computed **using** the Label Match **algorithm** detailed in. Section 4.  
Step 3 - Contextual **Alignment**: The similarity matrix ...  
[ieeexplore.ieee.org/iel5/10810/34089/01623879.pdf?isnumber=34089&prod=CNF&anumber=1623879&arSt=84&ared...K...](#) - [Similar pages](#)

9. [\[PDF\] SAMBO Results for the Ontology Alignment Evaluation Initiative 2007](#)

File Format: PDF/Adobe Acrobat - [View as HTML](#)

This **article** describes a system for ontology **alignment**, SAMBO, and .... matching **algorithms**, n-gram and **edit distance**, and a linguistic **algorithm**. ...  
[www.dit.univr.it/~p2p/OM-2007/12-o-SAMBO.pdf](#) - [Similar pages](#)  
by H Tan - [Cited by 2](#) - [Related articles](#) - [All 3 versions](#)

10. [\[PDF\] A Comparison of Alignment Models for Statistical Machine Translation](#)

File Format: PDF/Adobe Acrobat - [View as HTML](#)

**source** and target words **using** an **alignment** nmpling .... known an efficient **algorithm** to **compute** the Viterbi. **alignment** for the Models 3 and 4. ...  
[acl.ldc.upenn.edu/C/C00/C00-2163.pdf](#) - [Similar pages](#)  
by FJ Och - 2000 - [Cited by 94](#) - [Related articles](#) - [All 4 versions](#)



compare source article using alignment algorithm to compute edit distance

Search

[Search within results](#) | [Language Tools](#) | [Search Tips](#) | [Dissatisfied? Help us improve](#) | [Try Google Experimental](#)

[Google Home](#) - [Advertising Programs](#) - [Business Solutions](#) - [Privacy](#) - [About Google](#)